In the Claims

Please amend the claims as follows:

- 1-23. (Cancelled)
- 24. (Currently amended) The method of claim 23, A method for identifying elite event MS-B2 in a transgenic Brassica plant, or cell or tissue thereof, or transgenic Brassica plant material, said method comprising amplifying a DNA fragment of between 100 and 300 nucleotides 160 and 200 by from a nucleic acid present in said transgenic Brassica plant, or cell or tissue thereof, or transgenic Brassica plant material, using a polymerase chain reaction (PCR) with a first specific primer or probe at least-two-primers, one of which comprises at least 16 consecutive nucleotides from the 5' flanking region of MS-B2, comprised in hybridizes to bases 1-234 of SEQ ID NO:8, or the complement thereof, or from the 3' flanking region of MS-B2, comprised in to bases 194-416 of SEQ ID NO:10, or the complement thereof; of MS-B2, and a second specific primer or probe the other of which comprises at least 16 consecutive nucleotides from the foreign DNA in MS-B2, or the complement thereof, said foreign DNA corresponding to hybridizes to a sequence within SEQ ID NO:1; and thus identifying a Brassica plant, or cell or tissue thereof, or transgenic plant material comprising elite event MS-B2, if said genomic DNA amplifies the DNA fragment using PCR with the primers and detecting said amplified DNA fragment on an agarose gel.
- 25. (Currently amended) The method of claim 24, wherein one of said second specific primer or probe primers hybridizes to a sequence within SEQ ID NO:1 and comprises the sequence of SEQ ID NO: 12.
- 26. (Currently amended) The method of claim 24, wherein one of said first specific primer or probe comprises at least 16 consecutive nucleotides from the 3' flanking region of MS-B2, comprised in primers hybridizes to bases 194-416 of SEQ ID NO:10, or the complement thereof and comprises the sequence of SEQ ID NO:11.
 - 27-29. (Cancelled)
- 30. (Currently amended) The kit of Claim 29, which further comprises at least A kit for identifying elite event MS-B2 in a transgenic Brassica plant, or cell or tissue thereof, or transgenic Brasscia plant material, said kit comprising at least a first PCR primer or probe and a second PCR primer or probe, wherein the first PCR primer or probe comprises at least 16 consecutive nucleotides from the 5' flanking region of MS-B2, comprised in bases 1-234 of SEQ

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